

Jessica S Thomas

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

34
citations

1937632

4
h-index

1872665

6
g-index

10
all docs

10
docs citations

10
times ranked

64
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Molecular Signatures of <i>KRAS</i> -Mutated Lung Adenocarcinoma: Analysis of Concomitant <i>EGFR</i> , <i>ALK</i> , <i>STK11</i> , and PD-L1 Status. <i>BMC Clinical Pathology</i> , 2022, 15, 2632010X2211020. | 1.7 | 2 |
| 2 | Identifying possible germline variants from tumor-only sequencing of hematological malignancies. <i>Leukemia and Lymphoma</i> , 2021, 62, 482-485. | 1.3 | 0 |
| 3 | Pleomorphic giant cell carcinoma of prostate: Rare tumor with unique clinicopathological, immunohistochemical, and molecular features. <i>Annals of Diagnostic Pathology</i> , 2021, 52, 151719. | 1.3 | 9 |
| 4 | Pathology Trainee Redeployment and Education During the COVID-19 Pandemic: An Institutional Experience. <i>Academic Pathology</i> , 2020, 7, 2374289520953548. | 1.1 | 11 |
| 5 | Development and validation of Houston Methodist Variant Viewer version 3: updates to our application for interpretation of next-generation sequencing data. <i>JAMIA Open</i> , 2020, 3, 299-305. | 2.0 | 0 |
| 6 | A BRAF V600E Mutation in RET-Negative Medullary Thyroid Cancer. <i>Case Reports in Endocrinology</i> , 2020, 2020, 1-4. | 0.4 | 2 |
| 7 | Updates to a Lab-Developed Bioinformatics Pipeline and Application for Interpretation of Clinical Next-Generation Sequencing Panels. <i>American Journal of Clinical Pathology</i> , 2019, 152, S9-S10. | 0.7 | 0 |
| 8 | Mutational Landscape of Intermediate-Risk and Poor-Risk Acute Myeloid Leukemia (AML) Adult Patients by Next-Generation Sequencing (NGS): One Academic Center's Experience. <i>American Journal of Clinical Pathology</i> , 2018, 150, S154-S154. | 0.7 | 0 |
| 9 | Clinical Utility of Reflex Ordered Testing for Molecular Biomarkers in Lung Cancer. <i>American Journal of Clinical Pathology</i> , 2018, 150, S143-S143. | 0.7 | 5 |
| 10 | Houston Methodist Variant Viewer: An Application to Support Clinical Laboratory Interpretation of Next-generation Sequencing Data for Cancer. <i>Journal of Pathology Informatics</i> , 2017, 8, 44. | 1.7 | 5 |